OCT 2 1 2005

SEQUENCE LISTING

0> Ish-Horowicz, David
 Henrique , Domingos Manuel Pinto
 Lewis, Julian Hart
 Artavanis Tsakonas, Spyridon
 Gray, Grace

<120> ANTIBODIES TO VERTEBRATE DELTA PROTEINS

AND FRAGMENTS

- <130> 7326-122-999
- <140> 09/783,931
- <141> 2001-02-15
- <150> 08/981,392
- <151> 1997-12-22
- <150> PCT/US96/11178
- <151> 1996-06-28
- <150> 60/000,589
- <151> 1995-06-28
- <160> 95
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Pro Pro Cys Thr Tyr Gly Ser Ala Ile Thr Pro Val Leu Gly Ala Asn
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Ser Phe Ser Val Pro Asp Gly Ala Gly Gly Ala Asp Pro Ala Phe Ser
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Asn Pro Ile Arg Phe Pro Phe Gly Phe Thr Trp Pro Gly Thr Phe Ser
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Pro Trp Ile Ala Val Cys Ala Gly Ile Val Leu Val Leu Met Leu Leu
                550
                                      555
Leu Gly Cys Ala Ala Val Val Cys Val Arg Val Arg Val Gln Lys
                                  570
               565
Arg Arg His Gln Pro Glu Ala Cys Arg Gly Glu Ser Lys Thr Met Asn
                                                  590
           580
                              585
Asn Leu Ala Asn Cys Gln Arg Glu Lys Asp Ile Ser Val Ser Phe Ile
                           600
                                              605
Gly Thr Thr Gln Ile Lys Asn Thr Asn Lys Lys Ile Asp Phe Leu Ser
                      615
                                          620
Glu Ser Asn Asn Glu Lys Asn Gly Tyr Lys Pro Arg Tyr Pro Ser Val
                   630 635
Asp Tyr Asn Leu Val His Glu Leu Lys Asn Glu Asp Ser Pro Lys Glu
               645
                                  650
Glu Arg Ser Lys Cys Glu Ala Lys Cys Ser Ser Asn Asp Ser Asp Ser
                              665
                                                 670
Glu Asp Val Asn Ser Val His Ser Lys Arg Asp Ser Ser Glu Arg Arg
                                               685
                           680
Arg Pro Asp Ser Ala Tyr Ser Thr Ser Lys Asp Thr Lys Tyr Gln Ser
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Val Tyr Val Ile Ser Asp Glu Lys Asp Glu Cys Ile Ile Ala Thr Glu
705
                   710
                                       715
Val
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<213> Drosophila
<400> 6
Met His Trp Ile Lys Cys Leu Leu Thr Ala Phe Ile Cys Phe Thr Val
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Ile Val Gln Val His Ser Ser Gly Ser Phe Glu Leu Arg Leu Lys Tyr
                                25
Phe Ser Asn Asp His Gly Arg Asp Asn Glu Gly Arg Cys Cys Ser Gly
                            40
Glu Ser Asp Gly Ala Thr Gly Lys Cys Leu Gly Ser Cys Lys Thr Arg
                        55
Phe Arg Leu Cys Leu Lys His Tyr Gln Ala Thr Ile Asp Thr Thr Ser
                   70
                                        75
Gln Cys Thr Tyr Gly Asp Val Ile Thr Pro Ile Leu Gly Glu Asn Ser
Val Asn Leu Thr Asp Ala Gln Arg Phe Gln Asn Lys Gly Phe Thr Asn
                                105
Pro Ile Gln Phe Pro Phe Ser Phe Ser Trp Pro Gly Thr Phe Ser Leu
                            120
Ile Val Glu Ala Trp His Asp Thr Asn Asn Ser Gly Asn Ala Arg Thr
                       135
                                            140
Asn Lys Leu Leu Ile Gln Arg Leu Leu Val Gln Gln Val Leu Glu Val
                   150
                                        155
Ser Ser Glu Trp Lys Thr Asn Lys Ser Glu Ser Gln Tyr Thr Ser Leu
               165
                                    170
Glu Tyr Asp Phe Arg Val Thr Cys Asp Leu Asn Tyr Tyr Gly Ser Gly
           180
                               185
Cys Ala Lys Phe Cys Arg Pro Arg Asp Asp Ser Phe Gly His Ser Thr
                            200
Cys Ser Glu Thr Gly Glu Ile Ile Cys Leu Thr Gly Trp Gln Gly Asp
                       215
                                            220
Tyr Cys His Ile Pro Lys Cys Ala Lys Gly Cys Glu His Gly His Cys
                   230
                                        235
Asp Lys Pro Asn Gln Cys Val Cys Gln Leu Gly Trp Lys Gly Ala Leu
                                    250
               245
Cys Asn Glu Cys Val Leu Glu Pro Asn Cys Ile His Gly Thr Cys Asn
           260
                                265
Lys Pro Trp Thr Cys Ile Cys Asn Glu Gly Trp Gly Gly Leu Tyr Cys
                            280
Asn Gln Asp Leu Asn Tyr Cys Thr Asn His Arg Pro Cys Lys Asn Gly
                        295
                                            300
Gly Thr Cys Phe Asn Thr Gly Glu Gly Leu Tyr Thr Cys Lys Cys Ala
                   310
                                        315
Pro Gly Tyr Ser Gly Asp Asp Cys Glu Asn Glu Ile Tyr Ser Cys Asp
               325
                                    330
Ala Asp Val Asn Pro Cys Gln Asn Gly Gly Thr Cys Ile Asp Glu Pro
           340
                                345
His Thr Lys Thr Gly Tyr Lys Cys His Cys Arg Asn Gly Trp Ser Gly
                            360
                                                365
Lys Met Cys Glu Glu Lys Val Leu Thr Cys Ser Asp Lys Pro Cys His
                       375
                                            380
Gln Gly Ile Cys Arg Asn Val Arg Pro Gly Leu Gly Ser Lys Gly Gln
                   390
                                        395
Gly Tyr Gln Cys Glu Cys Pro Ile Gly Tyr Ser Gly Pro Asn Cys Asp
                405
                                    410
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<210> 6

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Leu Gln Leu Asp Asn Cys Ser Pro Asn Pro Cys Ile Asn Gly Gly Ser
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Cys Gln Pro Ser Gly Lys Cys Ile Cys Pro Ser Gly Phe Ser Gly Thr
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                            440
                                                445
Arg Cys Glu Thr Asn Ile Asp Asp Cys Leu Gly His Gln Cys Glu Asn
                        455
Gly Gly Thr Cys Ile Asp Met Val Asn Gln Tyr Arg Cys Gln Cys Val
                    470
                                        475
Pro Gly Phe His Gly Thr His Cys Ser Ser Lys Val Asp Leu Cys Leu
                485
                                    490
Ile Arg Pro Cys Ala Asn Gly Gly Thr Cys Leu Asn Leu Asn Asn Asp
            500
                                505
Tyr Gln Cys Thr Cys Arg Ala Gly Phe Thr Gly Lys Asp Cys Ser Val
       515
                            520
                                               525
Asp Ile Asp Glu Cys Ser Ser Gly Pro Cys His Asn Gly Gly Thr Cys
                        535
Met Asn Arg Val Asn Ser Phe Glu Cys Val Cys Ala Asn Gly Phe Arg
                    550
                                        555
Gly Lys Gln Cys Asp Glu Glu Ser Tyr Asp Ser Val Thr Phe Asp Ala
               565
                                   570
His Gln Tyr Gly Ala Thr Thr Gln Ala Arg Ala Asp Gly Leu Ala Asn
            580
                                585
Ala Gln Val Val Leu Ile Ala Val Phe Ser Val Ala Met Pro Leu Val
                            600
Ala Val Ile Ala Ala Cys Val Val Phe Cys Met Lys Arg Lys Arg Lys
                        615
Arg Ala Gln Glu Lys Asp Asn Ala Glu Ala Arg Lys Gln Asn Glu Gln
                    630
                                        635
Asn Ala Val Ala Thr Met His His Asn Gly Ser Ala Val Gly Val Ala
                645
                                    650
Leu Ala Ser Ala Ser Met Gly Gly Lys Thr Gly Ser Asn Ser Gly Leu
                                665
Thr Phe Asp Gly Gly Asn Pro Asn Ile Ile Lys Asn Thr Trp Asp Lys
                            680
Ser Val Asn Asn Ile Cys Ala Ser Ala Ala Ala Ala Ala Ala Ala Ala
                       695
Ala Ala Asp Glu Cys Leu Met Tyr Gly Gly Tyr Val Ala Ser Val
                    710
                                        715
Ala Asp Asn Asn Ala Asn Ser Asp Phe Cys Val Ala Pro Leu Gln
               725
                                    730
Arg Ala Lys Ser Gln Lys Gln Leu Asn Thr Asp Pro Thr Leu Met His
                                745
Arg Gly Ser Pro Ala Gly Thr Ser Ala Lys Gly Ala Ser Gly Gly
                            760
Pro Gly Ala Ala Glu Gly Lys Arg Ile Ser Val Leu Gly Glu Gly Ser
                        775
                                            780
Tyr Cys Ser Gln Arg Trp Pro Ser Leu Ala Ala Gly Val Ala Gly
                   790
                                        795
Asp Leu Phe Ile Gln Leu Met Ala Ala Ala Ser Val Ala Gly Thr Asp
               805
                                   810
                                                        815
Gly Thr Ala Gln Gln Arg Ser Val Val Cys Gly Thr Pro His Met
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Val Gln Cys Ala Val Thr Tyr Tyr Asn Thr Thr Phe Cys Thr Thr Phe

<210> 7 <211> 46

<212> PRT

<213> Drosophila

<400> 7

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Cys Arg Pro Arg Asp Asp Gln Phe Gly His Tyr Ala Cys Gly Ser Glu
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Gly Gln Lys Leu Cys Leu Asn Gly Trp Gln Gly Val Asn Cys
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<211> 45
<212> PRT
<213> Gallus gallus
<400> 8
Val Thr Cys Ala Glu His Tyr Tyr Gly Phe Gly Cys Asn Lys Phe Cys
Arg Pro Arg Asp Asp Phe Phe Thr His His Thr Cys Asp Gln Asn Gly
                                25
Asn Lys Thr Cys Leu Glu Gly Trp Thr Gly Pro Glu Cys
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<211> 43
<212> PRT
<213> Drosophila
<400> 9
Asn Leu Cys Ser Ser Asn Tyr His Gly Lys Arg Cys Asn Arg Tyr Cys
Ile Ala Asn Ala Lys Leu His Trp Glu Cys Ser Thr His Gly Val Arg
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            20
Arg Cys Ser Ala Gly Trp Ser Gly Glu Asp Cys
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<211> 45
<212> PRT
<213> Drosophila
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Val Thr Cys Ala Arg Asn Tyr Phe Gly Asn Arg Cys Glu Asn Phe Cys
Asp Ala His Leu Ala Lys Ala Ala Arg Lys Arg Cys Asp Ala Met Gly
            20
                                 2.5
Arg Leu Arg Cys Asp Ile Gly Trp Met Gly Pro His Cys
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<210> 11
<211> 2692
<212> DNA
<213> mouse
<220>
<221> CDS
<222> (31)...(2199)
<223> Mouse Delta (M-Delta-1) gene
<400> 11
ctgcaggaat tcsmycgcat gctcccggcc gcc atg ggc cgt cgg agc gcg cta 54
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Ala Met Gly Arg Arg Ser Ala Leu 1 5

			tct Ser										102
			ctg Leu 30										150
	-	_	tgc Cys	_	 			_		_	-	-	198
		_	gta Val	-	_			_	_	_			246
			acc Thr		 _	-	-	_			_		294
			ctg Leu										342
			ttc Phe 110										390
			gcc Ala										438
			aga Arg										486
			gaa Glu										534
			tct Ser										582
			gtg Val 190										630
			gac Asp										678
	_	_	act Thr	-		_	_			_	_	-	726
			gac Asp										774

235		240	245		
	cgc tac tgc gat Arg Tyr Cys Asp 255		-		
	tgc cag caa ccc Cys Gln Gln Pro 270				
	ttc tgc aac caa Phe Cys Asn Gln 285	-	_		•
	aat gga gcc acc Asn Gly Ala Thr 300	-			
-	tgc cga cct ggg Cys Arg Pro Gly		_		
, , ,	tgt gct cct agc Cys Ala Pro Ser 335		,,,,,	, ,	•
	gac agc ttc tct Asp Ser Phe Ser 350				7
	gag ctg agc gcc Glu Leu Ser Ala 365	-		-	
	cga tgt tca gat Arg Cys Ser Asp 380				
-	ggc ttc tct ggc Gly Phe Ser Gly	_			
	tcc cct tgt tct Ser Pro Cys Ser 415				
	ctg tgc cgg tgc Leu Cys Arg Cys 430				5
	gtg gat gac tgt Val Asp Asp Cys 445				
	gac agt gtg aac Asp Ser Val Asn 460		Cys Thr Cys		
	aag aac tgc agc Lys Asn Cys Ser				

											ggc Gly 500					1542
											tgc Cys					1590
							_			_	ctc Leu	_	-			1638
											gcc Ala					1686
			_		_	_	_	_		_	gct Ala	-			-	1734
_	_		_	-		_			_		cca Pro 580		-		_	1782
				-		_				_	aat Asn	_	_	_		1830
											cag Gln					1878
	_	_		_				-			gcc Ala		_	-	-	1926
	_	_	_					_			ctc Leu	-	_	-		1974
											agc Ser 660					2022
_	_	_		_	_	_	_		_		aga Arg					2070
											cca Pro					2118
											tat Tyr					2166
					gtt Val						taaq	gatg	gaa q	gcgat	tgtggc	2219

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aaaattccca tttctcttaa ataaaattcc aaggatatag ccccgatgaa tgctgctgag 2279
agaggaaggg agaggaaacc cagggactgc tgctgagaac caggttcagg cgaacgtggt 2339
teteteagag ttageagagg egecegaeae tgeeageeta ggetttgget geegetggae 2399
tgcctgctgg ttgttcccat tgcactatgg acagttgctt tgaagagtat atatttaaat 2459
ggacgagtga cttgattcat ataggaagca cgcactgccc acacgtctat cttggattac 2519
tatgagccag tettteettg aactagaaac acaactgeet ttattgteet ttttgataet 2579
gagatgtgtt ttttttttt cctagacggg aaaaagaaaa cgtgtgttat tttttttggg 2639
atttgtaaaa atattttca tgattatggg agagctccca acgcgttgga ggt
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Cys Gln Val Trp Ser Ser Gly Val Phe Glu Leu Lys Leu Gln Glu Phe
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                                                    30
Val Asn Lys Lys Gly Leu Leu Gly Asn Arg Asn Cys Cys Arg Gly Gly
                            40
Ser Gly Pro Pro Cys Ala Cys Arg Thr Phe Phe Arg Val Cys Leu Lys
His Tyr Gln Ala Ser Val Ser Pro Glu Pro Pro Cys Thr Tyr Gly Ser
                    70
                                        75
Ala Val Thr Pro Val Leu Gly Val Asp Ser Phe Ser Leu Pro Asp Gly
                                    90
Ala Gly Ile Asp Pro Ala Phe Ser Asn Pro Ile Arg Phe Pro Phe Gly
                                105
Phe Thr Trp Pro Gly Thr Phe Ser Leu Ile Ile Glu Ala Leu His Thr
                            120
Asp Ser Pro Asp Asp Leu Ala Thr Glu Asn Pro Glu Arg Leu Ile Ser
                        135
                                            140
Arg Leu Thr Thr Gln Arg His Leu Thr Val Gly Glu Glu Trp Ser Gln
                    150
                                        155
Asp Leu His Ser Ser Gly Arg Thr Asp Leu Arg Tyr Ser Tyr Arg Phe
                165
                                    170
Val Cys Asp Glu His Tyr Tyr Gly Glu Gly Cys Ser Val Phe Cys Arg
                                185
            180
Pro Arg Asp Asp Ala Phe Gly His Phe Thr Cys Gly Asp Arg Gly Glu
                            200
                                                205
Lys Met Cys Asp Pro Gly Trp Lys Gly Gln Tyr Cys Thr Asp Pro Ile
                        215
                                            220
Cys Leu Pro Gly Cys Asp Asp Gln His Gly Tyr Cys Asp Lys Pro Gly
                    230
                                        235
Glu Cys Lys Cys Arg Val Gly Trp Gln Gly Arg Tyr Cys Asp Glu Cys
                                    250
                245
Ile Arg Tyr Pro Gly Cys Val His Gly Thr Cys Gln Gln Pro Trp Gln
                                265
            260
Cys Asn Cys Gln Glu Gly Trp Gly Gly Leu Phe Cys Asn Gln Asp Leu
        275
                            280
                                                285
Asn Tyr Cys Thr His His Lys Pro Cys Arg Asn Gly Ala Thr Cys Thr
                                            300
                        295
Asn Thr Gly Gln Gly Ser Tyr Thr Cys Ser Cys Arg Pro Gly Tyr Thr
                    310
                                        315
Gly Ala Asn Cys Glu Leu Glu Val Asp Glu Cys Ala Pro Ser Pro Cys
                                    330
                325
                                                        335
Lys Asn Gly Ala Ser Cys Thr Asp Leu Glu Asp Ser Phe Ser Cys Thr
            340
                                345
                                                    350
Cys Pro Pro Gly Phe Tyr Gly Lys Val Cys Glu Leu Ser Ala Met Thr
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365

360

355

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Cys Ala Asp Gly Pro Cys Phe Asn Gly Gly Arg Cys Ser Asp Asn Pro
    370
                        375
                                            380
Asp Gly Gly Tyr Thr Cys His Cys Pro Leu Gly Phe Ser Gly Phe Asn
                    390
                                        395
Cys Glu Lys Lys Met Asp Leu Cys Gly Ser Ser Pro Cys Ser Asn Gly
                                    410
Ala Lys Cys Val Asp Leu Gly Asn Ser Tyr Leu Cys Arg Cys Gln Ala
                                425
Gly Phe Ser Gly Arg Tyr Cys Glu Asp Asn Val Asp Asp Cys Ala Ser
        435
                            440
                                                445
Ser Pro Cys Ala Asn Gly Gly Thr Cys Arg Asp Ser Val Asn Asp Phe
                        455
                                            460
Ser Cys Thr Cys Pro Pro Gly Tyr Thr Gly Lys Asn Cys Ser Ala Pro
                    470
                                        475
Val Ser Arg Cys Glu His Ala Pro Cys His Asn Gly Ala Thr Cys His
                                    490
                                                        495
                485
Gln Arg Gly Gln Arg Tyr Met Cys Glu Cys Ala Gln Gly Tyr Gly Gly
            500
                                505
                                                    510
Pro Asn Cys Gln Phe Leu Leu Pro Glu Pro Pro Pro Gly Pro Met Val
        515
                            520
                                                525
Val Asp Leu Ser Glu Arg His Met Glu Ser Gln Gly Pro Phe Pro
                        535
                                            540
Trp Val Ala Val Cys Ala Gly Val Val Leu Val Leu Leu Leu Leu
                    550
                                        555
Gly Cys Ala Ala Val Val Cys Val Arg Leu Lys Leu Gln Lys His
                565
                                    570
                                                        575
Gln Pro Pro Glu Pro Cys Gly Glu Thr Glu Thr Met Asn Asn
            580
                                585
Leu Ala Asn Cys Gln Arg Glu Lys Asp Val Ser Val Ser Ile Ile Gly
                            600
                                                605
        595
Ala Thr Gln Ile Lys Asn Thr Asn Lys Lys Ala Asp Phe His Gly Asp
                        615
                                            620
His Gly Ala Glu Lys Ser Ser Phe Lys Val Arg Tyr Pro Thr Val Asp
                    630
                                        635
Tyr Asn Leu Val Arg Asp Leu Lys Gly Asp Glu Ala Thr Val Arg Asp
                645
                                    650
                                                        655
Thr His Ser Lys Arg Asp Thr Lys Cys Gln Ser Gln Ser Leu Gln Glu
            660
                                665
                                                    670
Lys Arg Arg Ser Pro Gln His Leu Gly Val Gly Arg Phe Leu Thr Glu
                            680
                                                685
Asn Arg Pro Glu Ser Val Tyr Ser Thr Ser Lys Asp Thr Lys Tyr Gln
                        695
                                            700
Ser Val Tyr Val Leu Ser Ala Glu Lys Asp Glu Cys Val Ile Ala Thr
                    710
                                        715
Glu Val
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Asn Cys Cys Arg Gly Gly Cys Cys Thr Phe Phe Arg Val Cys Leu
                            40
Lys His Tyr Gln Ala Ser Val Ser Pro Glu Pro Pro Cys Thr Tyr Gly
                        55
Ser Ala Thr Pro Val Leu Gly Ser Phe Ser Pro Asp Gly Ala Gly Asp
                    70
                                        75
                                                             80
Pro Ala Phe Ser Asn Pro Ile Arg Phe Pro Phe Gly Phe Thr Trp Pro
                                    90
                                                         95
Gly Thr Phe Ser Leu Ile Ile Glu Ala Leu His Thr Asp Ser Pro Asp
            100
                                105
Asp Leu Thr Glu Asn Pro Glu Arg Leu Ile Ser Arg Leu Thr Gln Arg
        115
                            120
His Leu Val Gly Glu Glu Trp Ser Gln Asp Leu His Ser Ser Gly Arg
                        135
                                            140
Thr Asp Leu Tyr Ser Tyr Arg Phe Val Cys Asp Glu His Tyr Tyr Gly
                                        155
                    150
Glu Gly Cys Ser Val Phe Cys Arg Pro Arg Asp Asp Phe Gly His Phe
                                    170
                165
Thr Cys Gly Arg Gly Glu Lys Cys Pro Gly Trp Lys Gly Gln Tyr Cys
                                185
                                                     190
            180
Thr Pro Ile Cys Leu Pro Gly Cys Asp Gln His Gly Cys Asp Lys Pro
                            200
                                                 205
        195
Gly Glu Cys Lys Cys Arg Val Gly Trp Gln Gly Arg Tyr Cys Asp Glu
                                            220
                        215
Cys Ile Arg Tyr Pro Gly Cys Val His Gly Thr Cys Gln Gln Pro Trp
                    230
                                        235
Gln Cys Asn Cys Gln Glu Gly Trp Gly Gly Leu Phe Cys Asn Gln Asp
                245
                                    250
Leu Asn Tyr Cys Thr His His Lys Pro Cys Asn Gly Ala Thr Cys Thr
                                265
            260
Asn Thr Gly Gln Gly Ser Tyr Thr Cys Ser Cys Arg Pro Gly Tyr Thr
                            280
                                                285
Gly Cys Glu Glu Cys Pro Cys Lys Asn Gly Ser Cys Thr Asp Leu
                                             300
                        295
Glu Ser Ser Cys Thr Cys Pro Pro Gly Phe Tyr Gly Lys Cys Glu Leu
                    310
                                        315
Ser Ala Met Thr Cys Ala Asp Gly Pro Cys Phe Asn Gly Gly Arg Cys
                325
                                    330
Asp Asn Pro Asp Gly Gly Tyr Cys Cys Pro Leu Gly Ser Gly Phe Asn
            340
                                345
Cys Glu Lys Lys Asp Cys Ser Ser Pro Cys Asn Gly Ala Cys Val Asp
                            360
                                                 365
Leu Gly Asn Ser Tyr Cys Cys Gln Ala Gly Phe Gly Arg Cys Asp Asn
                        375
                                             380
Val Asp Asp Cys Ala Ser Pro Cys Asn Gly Gly Thr Cys Asp Val Asn
                    390
                                        395
Asp Ser Cys Thr Cys Pro Pro Gly Tyr Gly Lys Asn Cys Ser Pro Val
                                    410
Ser Arg Cys Glu His Pro Cys His Asn Gly Ala Thr Cys His Arg Arg
                                425
Tyr Cys Glu Cys Ala Gly Tyr Gly Gly Asn Cys Gln Phe Leu Leu Pro
                            440
Glu Pro Pro Gly Pro Val Asp Glu Glu Gln Phe Pro Trp Ala Val Cys
                        455
Ala Gly Leu Val Leu Leu Leu Gly Cys Ala Ala Val Val Cys Val
                    470
                                         475
Arg Leu Lys Gln Lys Pro Glu Cys Glu Thr Glu Thr Met Asn Asn Leu
                                     490
                485
Ala Asn Cys Gln Arg Glu Lys Asp Ser Ser Ile Gly Ala Thr Gln Ile
                                505
            500
Lys Asn Thr Asn Lys Lys Asp Phe His Asp Lys Lys Val Arg Tyr Pro
```

```
520
                                                 525
        515
Val Asp Tyr Asn Leu Val Leu Lys Val His Lys Lys Cys Ser Glu Glu
                        535
Lys Ala Leu Arg Lys Arg Pro Ser Val Tyr Ser Thr Ser Lys Asp Thr
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Lys Tyr Gln Ser Val Tyr Val Ser Glu Lys Asp Glu Cys Ile Ala Thr
                565
                                    570
Glu Val
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<211> 525
<212> DNA
<213> Homo sapiens
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ggcyacgtca gatcargaac accaacaaga aggcggactt ymcascgggg gaccasagcg 120
tccgacaaga atggmtttca aggcccgcta ccccagcgtg gactataact cgtgcaggac 180
ctcaagggtg acgacaccgc cgtcaggacg tcgcacagca agcgtgacac caagtgccag 240
tececagget ceteagggag gagaagggga eeeegaceae acteaggggk tgegtgetge 300
gggccgggct caggagggg tacctggggg gtgtcttcct ggaaccactg ctccgtttct 360
cttcccaaat gttctcatgc attcattgtg gattttctct attttccttt tagtggagaa 420
gcatctgaaa gaaaaaggcc ggactcgggc tgttcaactt caaaagacac caagtaccag 480
                                                                   525
teggtgtacg teatateega ggagaaggae gagtgegtea tegea
<210> 15
<211> 10
<212> PRT
<213> Artificial Sequence
<223> Predicted amino acid sequence of humna delta
<220>
<221> VARIANT
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<223> Xaa = Any Amino Acid
Tyr Asp Glu Xaa Pro Gly Glu Leu Pro Ala
<210> 16
<211> 44
<212> PRT
<213> Artificial Sequence
<223> Predicted amino acid sequence of humna delta
<220>
<221> VARIANT
<222> 11, 15, 23, 24, 28
<223> Xaa = Any Amino Acid
<400> 16
Glu Gly His Leu Ser Gln His His Arg Gly Xaa Val Arg Ser Xaa Thr
                                    10
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Pro Thr Arg Arg Thr Xaa Xaa Arg Gly Thr Xaa Ala Ser Asp Lys
            20
                                25
Asn Gly Phe Gln Gly Pro Leu Pro Gln Arg Gly Leu
                            40
<210> 17
<211> 118
<212> PRT
<213> Artificial Sequence
<220>
<223> Predicted amino acid sequence of humna delta
<220>
<221> VARIANT
<222> 41
<223> Xaa = Any Amino Acid
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Leu Val Gln Asp Leu Lys Gly Asp Asp Thr Ala Val Arg Thr Ser His
                 5
                                    10
Ser Lys Arg Asp Thr Lys Cys Gln Ser Pro Gly Ser Ser Gly Arg Arg
                                                     30
            20
                                25
Arg Gly Pro Arg Pro His Ser Gly Xaa Ala Cys Cys Gly Pro Gly Ser
                                                 4.5
                            40
Gly Gly Gly Thr Trp Gly Val Ser Ser Trp His Cys Ser Val Ser Leu
                        55
                                             60
Pro Lys Cys Ser His Ala Phe Ile Val Asp Phe Leu Tyr Phe Pro Phe
                                        75
                    70
Ser Gly Glu Ala Ser Glu Arg Lys Arg Pro Asp Ser Gly Cys Ser Thr
                85
                                    90
Ser Lys Asp Thr Lys Tyr Gln Ser Val Tyr Val Ile Ser Glu Glu Lys
            100
                                105
Asp Glu Cys Val Ile Ala
       115
<210> 18
<211> 173
<212> PRT
<213> Artificial Sequence
<220>
<223> Predicted amino acid sequence of human delta
<220>
<221> VARIANT
<222> 34, 35, 39, 44, 96
<223> Xaa = Any Amino Acid
<400> 18
Thr Met Asn Asn Leu Ala Asn Cys Gln Arg Glu Lys Asp Ile Ser Val
Ser Ile Ile Gly Ala Thr Ser Asp Gln Glu His Gln Glu Gly Gly
                                25
Leu Xaa Xaa Gly Gly Pro Xaa Pro Thr Arg Met Xaa Phe Lys Ala Arg
                            40
Tyr Pro Ser Val Asp Tyr Asn Ser Cys Arg Thr Ser Arg Val Thr Thr
                        55
```

Pro Pro Ser Gly Arg Arg Thr Ala Ser Val Thr Pro Ser Ala Ser Pro

```
70
                                         75
65
Gln Ala Pro Gln Gly Gly Glu Gly Asp Pro Asp His Thr Gln Gly Xaa
                                     90
Arg Ala Ala Gly Arg Ala Gln Glu Gly Val Pro Gly Gly Cys Leu Pro
                                 105
                                                     110
Gly Thr Thr Ala Pro Phe Leu Phe Pro Asn Val Leu Met His Ser Leu
        115
                             120
                                                 125
Trp Ile Phe Ser Ile Phe Leu Leu Val Glu Lys His Leu Lys Glu Lys
                        135
                                             140
Gly Arg Thr Arg Ala Val Gln Leu Gln Lys Thr Pro Ser Thr Ser Arg
                    150
                                         155
Cys Thr Ser Tyr Pro Arg Arg Arg Thr Ser Ala Ser Ser
                165
                                     170
<210> 19
<211> 60
<212> PRT
<213> Artificial Sequence
<220>
<223> Predicted amino acid sequence of human delta
<220>
<221> VARIANT
<222> 1, 19, 23, 32, 33, 36, 43
<223> Xaa = Any Amino Acid
<400> 19
Xaa Thr Trp Arg Thr Ala Ser Val Arg Arg Thr Ser Gln Ser Ala Ser
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Ser Gly Xaa Arg Gln Ile Xaa Asn Thr Asn Lys Lys Ala Asp Phe Xaa
                                 25
Xaa Gly Asp Xaa Ser Val Arg Gln Glu Trp Xaa Ser Arg Pro Ala Thr
                            40
Pro Ala Trp Thr Ile Thr Arg Ala Gly Pro Gln Gly
    50
                        55
<210> 20
<211> 11
<212> PRT
<213> Artificial Sequence
<223> Predicted amino acid sequence of human delta
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Arg His Arg Arg Gln Asp Val Ala Gln Gln Ala
<210> 21
<211> 61
<212> PRT
<213> Artificial Sequence
<220>
<223> Predicted amino acid sequence of human delta
<400> 21
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His Gln Val Pro Val Pro Arg Leu Leu Arg Glu Glu Lys Gly Thr Pro
                                    10
Thr Thr Leu Arg Gly Cys Val Leu Arg Ala Gly Leu Arg Arg Gly Tyr
                                25
Leu Gly Gly Val Phe Leu Glu Pro Leu Leu Arg Phe Ser Ser Gln Met
                            40
Phe Ser Cys Ile His Cys Gly Phe Ser Leu Phe Ser Phe
<210> 22
<211> 33
<212> PRT
<213> Artificial Sequence
<220>
<223> Predicted amino acid sequence of human delta
<400> 22
Lys Lys Lys Ala Gly Leu Gly Leu Phe Asn Phe Lys Lys Arg His Gln
                                    10
Val Pro Val Gly Val Arg His Ile Arg Gly Glu Gly Arg Val Arg His
            20
Arg
<210> 23
<211> 175
<212> PRT
<213> Artificial Sequence
<223> Predicted amino acid sequence of human delta
<220>
<221> VARIANT
<222> 25, 34, 35, 38, 97
<223> Xaa = Any Amino Acid
<400> 23
Thr Met Asn Asn Leu Ala Asn Cys Gln Arg Glu Lys Asp Ile Ser Val
Ser Ile Ile Gly Ala Thr Gly Ile Xaa Asn Thr Asn Lys Lys Ala Asp
                                25
Phe Xaa Xaa Gly Asp Xaa Ser Ser Asp Lys Asn Gly Phe Gln Lys Ala
Arg Tyr Pro Ser Val Asp Tyr Asn Leu Val Gln Asp Leu Lys Gly Asp
                        55
Asp Thr Ala Val Arg Thr Ser His Ser Lys Arg Asp Thr Lys Cys Gln
                    70
Ser Pro Gly Ser Ser Gly Arg Arg Gly Pro Arg Pro His Ser Gly
                                    90
Xaa Ala Cys Cys Gly Pro Gly Ser Gly Gly Gly Thr Trp Gly Val Ser
            100
                                105
                                                    110
Ser Trp Asn His Cys Ser Val Ser Leu Pro Lys Cys Ser His Ala Phe
                            120
                                                125
Ile Val Asp Phe Leu Tyr Phe Pro Phe Ser Gly Glu Ala Ser Glu Arg
                        135
                                            140
Lys Arg Pro Asp Ser Gly Cys Ser Thr Ser Lys Asp Thr Lys Tyr Gln
```

155

150

145

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<210> 24
<211> 2899
<212> DNA
<213> Artificial Sequence
<220>
<223> Consenses sequence of mouse delta and human delta
<220>
<221> misc feature
<222> 854, 973, 984, 1582, 1787, 1819, 1864, 1916, 1951, 2033,
2152, 2156, 2171, 2183, 2194, 2212, 2220, 2226, 2230, 2244,
2245, 2264, 2265, 2266, 2287
<223> n = A, T, C or G
<400> 24
gtccagcggt accatgggcc gtcggagcgc gctacccctt gccgtggtct ctgccctgct 60
gtgccaggtc tggagctccg gcgtatttga gctgaagctg caggagttcg tcaacaagaa 120
ggggctgctg gggaaccgca actgctgccg cgggggctct ggcccgcctt gcgcctgcag 180
gaccttettt egegtatgee teaaceacta eeaggeeage gtgteaeegg ageeaeeetg 240
cacctacggc agtgctgtca cgccagtgct gggtctcgac tccttcagcc tgcctsatkg 300
sgyasgsryc smccycgagg yckwcrgyaw csmyaagyyy gatatcgmmy tycggcttca 360
cctggccrgg yaccttctct ctgatyattg aagcyctcca yacagaytct ccygatgacc 420
tegeaacaga aaacceagaa agacteatea geegeetgre eacycagagg cacetsackg 480
tgggmgarga rtggtcycag gacctkcaca gyagcggccg cacrgacctc mrgtactcyt 540
accgsttygt gtgtgacgar cactactacg gagarggytg ctctgtkttc tgccgwccyc 600
gggaygaygc cttyggccac ttcacctgyg gggasmgwgg ggagaarrtg tgcraccctg 660
gctggaaagg scmgtactgc acwgasccra tctgyctgcc wggrtgtgat gascarcatg 720
gatwytgtga caaaccaggg gartgcaagt gcagagtkgg ctggcagggc cgstactgyg 780
atgagtgyat ccgytaycca ggytgtctcc atggcacctg ccagcarccc tggcagtgya 840
actgccagga aggntggggg ggccttttct gcaaccarga cctgaactac tgyacwcacc 900
ataagccstg cargaatgga gccacctgca acmaacacgg gccaggggga gctacacwtg 960
ktcyttggcc ggncykgggt ayanagggtg ccamctgyga agcttgggra ktrgaygagt 1020
tgttgmyccy agcccytggy aagaacggag sgagctksac ggaycttcgg agracagctw 1080
ctcytgyacc tgcccwcccg gcttctaygg caarrtctgt garytgagyg ccatgacctg 1140
tgcrgayggc ccttgcttya ayggrggwcg rtgytcagay arcccygayg gaggstacas 1200
ctgccrytgc cccktgggct wctcyggctt caactgtgag aagaaratkg ayywctgcrg 1260
ctettemecy tgttetaayg gtgccaaqtg tgtggacete ggyraykeyt acetgtgccg 1320
stgccaggcy ggcttctcsg ggaggyactg ygasgacaay gtggaygact gygcctcctc 1380
cccgtgygcm aaygggggca cctgccggga yrgygtgaac gacttgtcct gyacctgccc 1440
rectqqctac acgggcarga actgcagygc eccygycagc aggtgygagc aygcaccetg 1500
ccayaatggg gccacctgcc acsagagggg ccascgctay wtgtgygagt gygcccrrrg 1560
ctayggsggy cccaactgcc anttyctgct cccygaarcy gmccmccmgg scccayggtg 1620
gtggaamctc msykararrm aymtarragr gccrgggsgg gcccwtcccc tkggtggycg 1680
tgtgygccgg ggtsrtsctt gtcctcmtgc tgctgctggg ctgtgcygct gtggtggtct 1740
gcgtccggct gargctrcag aarcaccrgc cyccascyga mccctgnsgg ggrgagacrg 1800
araccatgaa caacctrgnc aaytgccagc gygagaagga crtytcwgty agcatcatyg 1860
gggnyacsca catcaagaac accaacaaga aggcggactt ycacggggac cayrgngccr 1920
asaagaryrg cttyaaggyc cgmtacccmr nkgtggacta taacctcgtk crrgacctca 1980
agggwgayga mrccrcsgtc agggayrcrc acagcaarcg tgacaccaag tgncagycmc 2040
agrgctcykg aggrgargag aaggggaycs ccgaccmaca ctyagggggt ggaggaagmw 2100
tcytgamaga aaaaggccrg astyygggyy trytcwactt tcaaargaca ancmangtac 2160
magtcggtgt nygtymtktc ygnagragga aggntgastg ygtyataggm rnytgaggtn 2220
gtaarntggn agcgatgtgg caannttccc atttctcksa aaknnnattc cmmggatata 2280
gcyccgntga atgctkctga gagaggaagg gagaggaaac ccagggactg ytkytcagaa 2340
ccaggttcag gcgaagctgg ttctctcaga gttagcagag gcgcccgaca ctgccagcct 2400
aggetttgge tgeegetgga etgeetgetg gttgtteeca ttgeactatg gaeagttget 2460
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Ser Val Tyr Val Ile Ser Glu Glu Lys Asp Glu Cys Val Ile Ala

170

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ttgaagagta tatatttaaa tggacgagtg acttgattca tatacgaagc acgcactgcc 2520
cacacqtcta tcttggatta ctatgagcca gtctttcctt gaactagaaa cacaactgcc 2580
cgtgtgttat ttttttggga tttgtaaaaa tatttttcat gatatctgta aagcttgagt 2700
attttgtgac gttcattttt ttataattta aattttggta aatatgtaca aaggcacttc 2760
gggtctatgt gactatattt ttttgtatat aaatgtattt atggaatatt gtgcaaatgt 2820
tatttgagtt ttttactgtt ttgttaatga agaaattcat tttaaaaaata tttttccaaa 2880
ataaatataa tgaactaca
                                                                 2899
<210> 25
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> sequence encoded by SEQ ID NO. 93 (degenerated
      oligo)
<400> 25
Glu Lys Asp Glu Cys Val Ile Ala
<210> 26
<211> 1981
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 559, 678, 689, 1287, 1492, 1524, 1569, 1621, 1656, 1738,
1857, 1861, 1876, 1888, 1899, 1917, 1925, 1931, 1935, 1942,
1943, 1952, 1953, 1954, 1968
<223> n = A, T, C or G
<400> 26
cattgggtac gggccccct cgaggtcgac ggtatcgata agcttgatat cgaattccgg 60
cttcacctgg ccgggcacct tctctctgat tattgaagct ctccacacag attctcctga 120
tgacctcgca acagaaaacc cagaaagact catcagccgc ctggccaccc agaggcacct 180
gacggtgggc gaggagtggt cccaggacct gcacagcagc ggccgcacgg acctcaagta 240
ctcctaccgc ttcgtgtgtg acgaacacta ctacggagag ggctgctccg ttttctgccg 300
teccegggae gatgeetteg gecaetteae etgtggggag egtggggaga aagtgtgeaa 360
ccctggctgg aaagggccct actgcacaga gccgatctgc ctgcctggat gtgatgagca 420
gcatggattt tgtgacaaac caggggaatg caagtgcaga gtgggctggc agggccggta 480
ctgtgacgag tgtatccgct atccaggctg tctccatggc acctgccagc agccctggca 540
gtgcaactgc caggaaggnt ggggggcct tttctgcaac caggacctga actactgcac 600
acaccataag ccctgcaaga atggagccac ctgcaacaaa cacgggccag ggggagctac 660
acttggtctt tggccggnct ggggtacana gggtgccacc tgcgaagctt ggggattgga 720
cgagttgttg accccagccc ttggtaagaa cggagggagc ttgacggatc ttcggagaac 780
agctactcct gtacctgccc acccggcttc tacggcaaaa tctgtgaatt gagtgccatg 840
acctgtgcgg acggcccttg ctttaacggg ggtcggtgct cagacagccc cgatggaggg 900
tacagctgcc gctgccccgt gggctactcc ggcttcaact gtgagaagaa aattgactac 960
tgcagctctt caccctgttc taatggtgcc aagtgtgtgg acctcggtga tgcctacctg 1020
tgccgctgcc aggccggctt ctcggggagg cactgtgacg acaacgtgga cgactgcgcc 1080
tecteeeegt gegeeaacgg gggeacetge egggatggeg tgaacgaett etectgeace 1140
tgcccgcctg gctacacggg caggaactgc agtgcccccg ccagcaggtg cgagcacgca 1200
ccctgccaca atggggccac ctgccacgag aggggccacc gctatttgtg cgagtgtgcc 1260
cgaagctacg ggggtcccaa ctgccanttc ctgctccccg aaactgcccc cccggcccca 1320
cggtggtgga aactccccta aaaaaaccta aaagggccgg ggggggccca tccccttggt 1380
ggacgtgtgc gccggggtca tccttgtcct catgctgctg ctgggctgtg ccgctgtggt 1440
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ggtctgcgtc cggctgaggc tgcagaagca ccggccccca gccgacccct gncggggga 1500
qacqqaqacc atqaacaacc tqgncaactg ccagcgtgag aaqqacatct caqtcaqcat 1560
categggne acgeagatea agaacaccaa caagaaggeg gaetteeacg gggaecacag 1620
ngccgacaag aatggcttca aggcccgcta cccagnggtg gactataacc tcqtqcaqqa 1680
cctcaagggt gacgacaccg ccgtcaggga cgcgcacagc aagcgtgaca ccaagtgnca 1740
gccccagggc tcctcagggg aggagaaggg gacccccgac ccacactcag ggggtggagg 1800
aagcatcttg aaagaaaaag gccggacttc gggcttgttc aactttcaaa agacaancaa 1860
ngtacaagtc ggtgtncgtc atttccgnag gaggaaggnt gactgcgtca taggaanttg 1920
aggtngtaaa ntggnagttg annttggaaa gnnntccccg gattccgntt tcaaagtttt 1980
<210> 27
<211> 31
<212> PRT
<213> Artificial Sequence
<220>
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<400> 27
His Trp Val Arg Ala Pro Leu Glu Val Asp Gly Ile Asp Lys Leu Asp
                 5
                                    10
Ile Glu Phe Arg Leu His Leu Ala Gly His Leu Leu Ser Asp Tyr
            20
                                25
<210> 28
<211> 7
<212> PRT
<213> Artificial Sequence
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<400> 28
Ser Ser Pro His Arg Phe Ser
<210> 29
<211> 45
<212> PRT
<213> Artificial Sequence
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<400> 29
Pro Arg Asn Arg Lys Pro Arg Lys Thr His Gln Pro Pro Gly His Pro
                                    10
Glu Ala Pro Asp Gly Gly Arg Gly Val Val Pro Gly Pro Ala Gln Gln
                                25
Arg Pro His Gly Pro Gln Val Leu Leu Pro Leu Arg Val
<210> 30
<211> 49
```

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<212> PRT
<213> Artificial Sequence
<220>
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<400> 30
Arg Thr Leu Leu Arg Arg Gly Leu Leu Arg Phe Pro Ser Pro Gly Arg
                 5
                                                         15
1
                                    10
Cys Leu Arg Pro Leu His Leu Trp Gly Ala Trp Gly Glu Ser Val Gln
            20
                                25
Pro Trp Leu Glu Arg Ala Leu Leu His Arg Ala Asp Leu Pro Ala Trp
                            40
Met
<210> 31
<211> 5
<212> PRT
<213> Artificial Sequence
<220>
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<400> 31
Ala Ala Trp Ile Leu
<210> 32
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<400> 32
Gln Thr Arg Gly Met Gln Val Gln Ser Gly Leu Ala Gly Pro Val Leu
<210> 33
<211> 40
<212> PRT
<213> Artificial Sequence
<220>
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<220>
<221> VARIANT
<222> 25
<223> Xaa = Any Amino Acid
<400> 33
```

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Arg Val Tyr Pro Leu Ser Arg Leu Ser Pro Trp His Leu Pro Ala Ala
 1
                 5
Leu Ala Val Gln Leu Pro Gly Arg Xaa Gly Gly Pro Phe Leu Gln Pro
Gly Pro Glu Leu Leu His Thr Pro
<210> 34
<211> 45
<212> PRT
<213> Artificial Sequence
<220>
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<220>
<221> VARIANT
<222> 27
<223> Xaa = Any Amino Acid
<400> 34
Ala Leu Gln Glu Trp Ser His Leu Gln Gln Thr Arg Ala Arg Gly Ser
                 5
                                    10
Tyr Thr Trp Ser Leu Ala Gly Leu Gly Tyr Xaa Gly Cys His Leu Arg
            20
                                25
Ser Leu Gly Ile Gly Arg Val Val Asp Pro Ser Pro Trp
<210> 35
<211> 196
<212> PRT
<213> Artificial Sequence
<220>
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<220>
<221> VARIANT
<222> 166, 179
<223> Xaa = Any Amino Acid
<400> 35
Glu Arg Arg Glu Leu Asp Gly Ser Ser Glu Asn Ser Tyr Ser Cys Thr
                                    10
Cys Pro Pro Gly Phe Tyr Gly Lys Ile Cys Glu Leu Ser Ala Met Thr
                                25
Cys Ala Asp Gly Pro Cys Phe Asn Gly Gly Arg Cys Ser Asp Pro Asp
                            40
Gly Gly Tyr Ser Cys Arg Cys Pro Val Gly Tyr Ser Gly Phe Asn Cys
                        55
Glu Lys Lys Ile Asp Tyr Cys Ser Ser Ser Pro Cys Ser Asn Gly Ala
                    70
                                        75
Lys Cys Val Asp Leu Gly Asp Ala Tyr Leu Cys Arg Gly Gln Ala Gly
                                    90
Phe Ser Gly Arg His Cys Asp Asp Asn Val Asp Asp Cys Ala Ser Ser
                                105
Pro Cys Ala Asn Gly Gly Thr Cys Arg Asp Gly Val Asn Asp Phe Ser
```

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120
        115
Cys Thr Cys Pro Pro Gly Tyr Thr Gly Arg Asn Cys Ser Ala Pro Ala
                        135
Ser Arg Cys Glu His Ala Pro Cys His Asn Gly Ala Thr Cys His Glu
                                         155
Arg Gly His Arg Tyr Xaa Cys Glu Cys Ala Arg Ser Tyr Gly Gly Pro
                165
                                     170
                                                         175
Asn Cys Xaa Phe Leu Leu Pro Glu Thr Ala Pro Pro Ala Pro Arg Trp
                                 185
Trp Lys Leu Pro
        195
<210> 36
<211> 65
<212> PRT
<213> Artificial Sequence
<220>
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<220>
<221> VARIANT
<222> 51
<223> Xaa = Any Amino Acid
<400> 36
Lys Asn Leu Lys Gly Pro Gly Gly Ala His Pro Leu Gly Gly Arg Val
                                    10
Arg Arg Gly His Pro Cys Pro His Ala Ala Gly Leu Cys Arg Cys
            20
                                25
Gly Gly Leu Arg Pro Ala Glu Ala Ala Glu Ala Pro Ala Pro Ser Arg
                            40
Pro Leu Xaa Gly Gly Asp Gly Asp His Glu Gln Pro Gly Gln Leu Pro
                        55
Ala
65
<210> 37
<211> 42
<212> PRT
<213> Artificial Sequence
<220>
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<220>
<221> VARIANT
<222> 28, 39
<223> Xaa = Any Amino Acid
<400> 37
Glu Gly His Leu Ser Gln His His Arg Gly His Ala Asp Gln Glu His
                                    10
Gln Glu Gly Gly Leu Pro Arg Gly Pro Gln Xaa Arg Gln Glu Trp
            20
                                25
Leu Gln Gly Pro Leu Pro Xaa Gly Gly Leu
```

```
<210> 38
<211> 7
<212> PRT
<213> Artificial Sequence
<220>
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<400> 38
Pro Arg Ala Gly Pro Gln Gly
                 5
 1
<210> 39
<211> 11
<212> PRT
<213> Artificial Sequence
<220>
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<400> 39
Arg His Arg Arg Gln Gly Arg Ala Gln Gln Ala
 1
<210> 40
<211> 57
<212> PRT
<213> Artificial Sequence
<220>
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<220>
<221> VARIANT
<222> 4, 43, 45, 50, 54
<223> Xaa = Any Amino Acid
<400> 40
His Gln Val Xaa Ala Pro Gly Leu Leu Arg Gly Gly Glu Gly Asp Pro
                 5
                                     10
Arg Pro Thr Leu Arg Gly Trp Arg Lys His Leu Glu Arg Lys Arg Pro
                                25
            20
                                                     30
Asp Phe Gly Leu Val Gln Leu Ser Lys Asp Xaa Gln Xaa Thr Ser Arg
       35
                            40
Cys Xaa Ser Phe Pro Xaa Glu Glu Gly
    50
<210> 41
<211> 8
<212> PRT
<213> Artificial Sequence
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<220>
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<220>
<221> VARIANT
<222> 5, 8
<223> Xaa = Any Amino Acid
<400> 41
Leu Arg His Arg Xaa Leu Arg Xaa
<210> 42
<211> 13
<212> PRT
<213> Artificial Sequence
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<220>
<221> VARIANT
<222> 1, 4, 5
<223> Xaa = Any Amino Acid
<400> 42
Xaa Trp Lys Xaa Xaa Pro Gly Phe Arg Phe Gln Ser Phe
<210> 43
<211> 276
<212> PRT
<213> Artificial Sequence
<220>
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<220>
<221> VARIANT
<222> 226, 230
<223> Xaa = Any Amino Acid
<400> 43
Ile Gly Tyr Gly Pro Pro Ser Arg Ser Thr Val Ser Ile Ser Leu Ile
                                    10
Ser Asn Ser Gly Phe Thr Trp Pro Gly Thr Phe Ser Leu Ile Ile Glu
            20
                                25
                                                     30
Ala Leu His Thr Asp Ser Pro Asp Asp Leu Ala Thr Glu Asn Pro Glu
                            40
                                                 45
Arg Leu Ile Ser Arg Leu Ala Thr Gln Arg His Leu Thr Val Gly Glu
                        55
Glu Trp Ser Gln Asp Leu His Ser Ser Gly Arg Thr Asp Leu Lys Tyr
                    70
                                         75
Ser Tyr Arg Phe Val Cys Asp Glu His Tyr Tyr Gly Glu Gly Cys Ser
```

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85
                                    90
Val Phe Cys Arg Pro Arg Asp Asp Ala Phe Gly His Phe Thr Cys Gly
            100
                                105
Glu Arg Gly Glu Lys Val Cys Asn Pro Gly Trp Lys Gly Pro Tyr Cys
       115
                            120
                                                125
Thr Glu Pro Ile Cys Leu Pro Gly Cys Asp Glu Gln His Gly Phe Cys
                       135
                                            140
Asp Lys Pro Gly Glu Cys Lys Cys Arg Val Gly Trp Gln Gly Arg Tyr
                    150
                                        155
Cys Asp Glu Cys Ile Arg Tyr Pro Gly Cys Leu His Gly Thr Cys Gln
                                    170
Gln Pro Trp Gln Cys Asn Cys Gln Glu Gly Trp Gly Gly Leu Phe Cys
                                185
Asn Gln Asp Leu Asn Tyr Cys Thr His His Lys Pro Cys Lys Asn Gly
                            200
Ala Thr Cys Asn Lys His Gly Pro Gly Gly Ala Thr Leu Gly Leu Trp
                        215
                                            220
Pro Xaa Trp Gly Thr Xaa Gly Ala Thr Cys Glu Ala Trp Gly Leu Asp
                    230
                                        235
Glu Leu Leu Thr Pro Ala Leu Gly Lys Asn Gly Gly Ser Leu Thr Asp
               245
                                    250
Leu Arg Arg Thr Ala Thr Pro Val Pro Ala His Pro Ala Ser Thr Ala
                                265
           260
Lys Ser Val Asn
       275
<210> 44
<211> 93
<212> PRT
<213> Artificial Sequence
<223> Deduced amino acid sequence using the three
     possible ORF of human Delta contigs
Pro Val Arg Thr Ala Leu Ala Leu Thr Gly Val Gly Ala Gln Thr Ala
                                    10
Pro Met Glu Gly Thr Ala Ala Ala Pro Trp Ala Thr Pro Ala Ser
            20
                                2.5
Thr Val Arg Arg Lys Leu Thr Thr Ala Ala Leu His Pro Val Leu Met
                            40
Val Pro Ser Val Trp Thr Ser Val Met Pro Thr Cys Ala Ala Arg
                        55
                                            60
Pro Ala Ser Arg Gly Gly Thr Val Thr Thr Thr Trp Thr Thr Ala Pro
                    70
                                        75
Pro Pro Arg Ala Pro Thr Gly Ala Pro Ala Gly Met Ala
               85
<210> 45
<211> 74
<212> PRT
<213> Artificial Sequence
```

<223> Deduced amino acid sequence using the three
 possible ORF of human Delta contigs

<220>

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<220>
<221> VARIANT
<222> 55
<223> Xaa = Any Amino Acid
<400> 45
Thr Thr Ser Pro Ala Pro Ala Arg Leu Ala Thr Arg Ala Gly Thr Ala
                                    10
Val Pro Pro Pro Ala Gly Ala Ser Thr His Pro Ala Thr Met Gly Pro
                                25
Pro Ala Thr Arg Gly Ala Thr Ala Ile Cys Ala Ser Val Pro Glu Ala
                            40
                                                45
Thr Gly Val Pro Thr Ala Xaa Ser Cys Pro Lys Leu Pro Pro Arg Pro
His Gly Gly Gly Asn Ser Pro Lys Lys Thr
<210> 46
<211> 187
<212> PRT
<213> Artificial Sequence
<220>
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<220>
<221> VARIANT
<222> 47, 58, 73, 101, 128, 167, 168, 181, 187
<223> Xaa = Any Amino Acid
<400> 46
Lys Gly Arg Gly Gly Pro Ile Pro Leu Val Asp Val Cys Ala Gly Val
                                    10
Ile Leu Val Leu Met Leu Leu Gly Cys Ala Ala Val Val Cys
            20
                                25
Val Arg Leu Arg Leu Gln Lys His Arg Pro Pro Ala Asp Pro Xaa Arg
        35
                            40
Gly Glu Thr Glu Thr Met Asn Asn Leu Xaa Asn Cys Gln Arg Glu Lys
                        55
Asp Ile Ser Val Ser Ile Ile Gly Xaa Thr Gln Ile Lys Asn Thr Asn
                    70
                                        75
Lys Lys Ala Asp Phe His Gly Asp His Ala Asp Lys Asn Gly Phe Lys
                85
                                    90
Ala Arg Tyr Pro Xaa Val Asp Tyr Asn Leu Val Gln Asp Leu Lys Gly
            100
                                105
Asp Asp Thr Ala Val Arg Asp Ala His Ser Lys Arg Asp Thr Lys Xaa
                            120
Gln Pro Gln Gly Ser Ser Gly Glu Glu Gly Thr Pro Asp Pro His Ser
                        135
Gly Gly Gly Ser Ile Leu Lys Glu Lys Gly Arg Thr Ser Gly Leu
                    150
                                        155
Phe Asn Phe Gln Lys Thr Xaa Xaa Val Gln Val Gly Val Arg His Phe
                165
                                    170
Arg Arg Arg Lys Xaa Asp Cys Val Ile Gly Xaa
<210> 47
```

<211> 20

```
<212> PRT
<213> Artificial Sequence
<220>
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<220>
<221> VARIANT
<222> 2, 4, 5, 7, 8, 11, 16
<223> Xaa = Any Amino Acid
Gly Xaa Lys Xaa Xaa Val Xaa Xaa Gly Lys Xaa Ser Pro Asp Ser Xaa
Phe Lys Val Phe
            20
<210> 48
<211> 12
<212> PRT
<213> Artificial Sequence
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
Leu Gly Thr Gly Pro Pro Arg Gly Arg Arg Tyr Arg
                 5
<210> 49
<211> 13
<212> PRT
<213> Artificial Sequence
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
Tyr Arg Ile Pro Ala Ser Pro Gly Arg Ala Pro Ser Leu
                 5
<210> 50
<211> 30
<212> PRT
<213> Artificial Sequence
<220>
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<400> 50
Leu Leu Lys Leu Ser Thr Gln Ile Leu Leu Met Thr Ser Gln Gln Lys
                 5
Thr Gln Lys Asp Ser Ser Ala Ala Trp Pro Pro Arg Gly Thr
```

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<210> 51
<211> 135
<212> PRT
<213> Artificial Sequence
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<220>
<221> VARIANT
<222> 126
<223> Xaa = Any Amino Acid
<400> 51
Arg Trp Ala Arg Ser Gly Pro Arg Thr Cys Thr Ala Ala Ala Arg
                                    10
Thr Ser Ser Thr Pro Thr Ala Ser Cys Val Thr Asn Thr Thr Glu
Arg Ala Ala Pro Phe Ser Ala Val Pro Gly Thr Met Pro Ser Ala Thr
                            40
Ser Pro Val Cys Ser Val Gly Arg Lys Cys Ala Thr Leu Ala Gly Lys
                        55
                                            60
Gly Pro Thr Ala Gln Ser Arg Ser Ala Cys Leu Asp Val Met Ser Ser
                    70
                                        75
Met Asp Phe Phe Val Thr Asn Gln Asn Ala Ser Ala Glu Trp Ala Gly
                85
                                    90
Arg Ala Gly Thr Val Thr Ser Val Ser Ala Ile Gln Ala Val Ser Met
            100
                                105
Ala Pro Ala Ser Ser Pro Gly Ser Ala Thr Ala Arg Lys Xaa Gly Gly
        115
                            120
                                                125
Ala Phe Ser Ala Thr Arg Thr
    130
                        135
<210> 52
<211> 46
<212> PRT
<213> Artificial Sequence
<220>
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<220>
<221> VARIANT
<222> 30, 33
<223> Xaa = Any Amino Acid
Thr Thr Ala His Thr Ile Ser Pro Ala Arg Met Glu Pro Pro Ala Thr
Asn Thr Gly Gln Gly Glu Leu His Leu Val Phe Gly Arg Xaa Gly Val
            20
                                25
Xaa Arg Val Pro Pro Ala Lys Leu Gly Asp Trp Thr Ser Cys
```

```
<210> 53
<211> 10
<212> PRT
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<220>
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<400> 53
Pro Gln Pro Leu Val Arg Thr Glu Gln Glu
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<210> 54
<211> 20
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<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<400> 54
Arg Ile Phe Gly Glu Gln Leu Leu Tyr Leu Pro Thr Arg Leu Leu
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                                   10
Arg Gln Asn Leu
<210> 55
<211> 12
<212> PRT
<213> Artificial Sequence
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<223> Deduced amino acid sequence using the three
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<400> 55
Ile Glu Cys His Asp Leu Cys Gly Arg Pro Leu Leu
<210> 56
<211> 25
<212> PRT
<213> Artificial Sequence
<220>
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<400> 56
Arg Gly Ser Val Leu Arg Gln Pro Arg Trp Arg Val Gln Leu Pro Leu
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Pro Arg Gly Leu Leu Arg Leu Gln Leu
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<210> 57

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<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Deduced amino acid sequence using the three
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<400> 57
Leu Leu Gln Leu Phe Thr Leu Phe
<210> 58
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
Trp Cys Gln Val Cys Gly Pro Arg
<210> 59
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
Cys Leu Pro Val Pro Leu Pro Gly Arg Leu Leu Gly Glu Ala Leu
                                    10
<210> 60
<211> 131
<212> PRT
<213> Artificial Sequence
<220>
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<220>
<221> VARIANT
<222> 76
<223> Xaa = Any Amino Acid
<400> 60
Arg Gln Arg Gly Arg Leu Arg Leu Pro Val Arg Gln Gly His Leu
                                    10
Pro Gly Trp Arg Glu Arg Leu Leu His Leu Pro Ala Trp Leu His
                                25
Gly Gln Glu Leu Gln Cys Pro Arg Gln Gln Val Arg Ala Arg Thr Leu
```

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40
                                                 45
        35
Pro Gln Trp Gly His Leu Pro Arg Glu Gly Pro Pro Leu Phe Val Arg
                        55
                                             60
Val Cys Pro Lys Leu Arg Gly Ser Gln Leu Pro Xaa Pro Ala Pro Arg
                    70
                                        75
Asn Cys Pro Pro Gly Pro Thr Val Val Glu Thr Pro Leu Lys Lys Pro
                85
                                    90
Lys Arg Ala Gly Gly Pro Ser Pro Trp Trp Thr Cys Ala Pro Gly
           100
                                105
                                                    110
Ser Ser Leu Ser Ser Cys Cys Cys Trp Ala Val Pro Leu Trp Trp Ser
                            120
Ala Ser Gly
    130
<210> 61
<211> 18
<212> PRT
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<223> Deduced amino acid sequence using the three
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<220>
<221> VARIANT
<222> 12
<223> Xaa = Any Amino Acid
<400> 61
Gly Cys Arg Ser Thr Gly Pro Gln Pro Thr Pro Xaa Gly Gly Arg Arg
Arg Pro
<210> 62
<211> 98
<212> PRT
<213> Artificial Sequence
<220>
<223> Deduced amino acid sequence using the three
     possible ORF of human Delta contigs
<220>
<221> VARIANT
<222> 4, 19, 36, 48, 75
<223> Xaa = Any Amino Acid
<400> 62
Thr Thr Trp Xaa Thr Ala Ser Val Arg Arg Thr Ser Gln Ser Ala Ser
                                    10
Ser Gly Xaa Arg Arg Ser Arg Thr Pro Thr Arg Arg Arg Thr Ser Thr
Gly Thr Thr Xaa Pro Thr Arg Met Ala Ser Arg Pro Ala Thr Gln Xaa
                            40
Trp Thr Ile Thr Ser Cys Arg Thr Ser Arg Val Thr Thr Pro Pro Ser
                        55
                                            60
Gly Thr Arg Thr Ala Ser Val Thr Pro Ser Xaa Ser Pro Arg Ala Pro
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Gln Gly Arg Arg Cys Pro Pro Thr His Thr Gln Gly Val Glu Glu
                85
                                    90
Ala Ser
<210> 63
<211> 33
<212> PRT
<213> Artificial Sequence
<220>
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<220>
<221> VARIANT
<222> 16, 17, 22, 26, 30
<223> Xaa = Any Amino Acid
<400> 63
Lys Lys Ala Gly Leu Arg Ala Cys Ser Thr Phe Lys Arg Gln Xaa
                                    10
Xaa Tyr Lys Ser Val Xaa Val Ile Ser Xaa Gly Gly Arg Xaa Thr Ala
            20
                                25
Ser
<210> 64
<211> 22
<212> PRT
<213> Artificial Sequence
<223> Deduced amino acid sequence using the three
      possible ORF of human Delta contigs
<220>
<221> VARIANT
<222> 2, 6, 8, 10, 13, 14, 19
<223> Xaa = Any Amino Acid
<400> 64
Glu Xaa Glu Val Val Xaa Trp Xaa Leu Xaa Leu Glu Xaa Xaa Pro Arg
                                    10
Ile Pro Xaa Ser Lys Phe
            20
<210> 65
<211> 192
<212> PRT
<213> Artificial Sequence
<220>
<223> Composite human delta (H-Delta-1) amino acid
     sequence
<400> 65
Gly Phe Thr Trp Pro Gly Thr Phe Ser Leu Ile Ile Glu Ala Leu His
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1
                 5
                                    10
Thr Asp Ser Pro Asp Asp Leu Ala Thr Glu Asn Pro Glu Arg Leu Ile
            20
                                25
Ser Arg Leu Ala Thr Gln Arg His Leu Thr Val Gly Glu Glu Trp Ser
                             40
Gln Asp Leu His Ser Ser Gly Arg Thr Asp Leu Lys Tyr Ser Tyr Arg
                        55
                                            60
Phe Val Cys Asp Glu His Tyr Tyr Gly Glu Gly Cys Ser Val Phe Cys
                    70
                                         75
Arg Pro Arg Asp Asp Ala Phe Gly His Phe Thr Cys Gly Glu Arg Gly
                85
                                    90
Glu Lys Val Cys Asn Pro Gly Trp Lys Gly Pro Tyr Cys Thr Glu Pro
                                105
Ile Cys Leu Pro Gly Cys Asp Glu Gln His Gly Phe Cys Asp Lys Pro
                            120
Gly Glu Cys Lys Cys Arg Val Gly Trp Gln Gly Arg Tyr Cys Asp Glu
                        135
                                            140
Cys Ile Arg Tyr Pro Gly Cys Leu His Gly Thr Cys Gln Gln Pro Trp
                    150
                                        155
Gln Cys Asn Cys Gln Glu Gly Trp Gly Gly Leu Phe Cys Asn Gln Asp
                165
                                    170
Leu Asn Tyr Cys Thr His His Lys Pro Cys Lys Asn Gly Ala Thr Cys
                                185
<210> 66
<211> 6
<212> PRT
<213> Artificial Sequence
<220>
<223> Composite human delta (H-Delta-1) amino acid
      sequence
<400> 66
Thr Asn Thr Gly Gln Gly
<210> 67
<211> 9
<212> PRT
<213> Artificial Sequence
<220>
<223> Composite human delta (H-Delta-1) amino acid
      sequence
<400> 67
Lys Asn Gly Gly Ser Leu Thr Asp Leu
<210> 68
<211> 157
<212> PRT
<213> Artificial Sequence
<220>
<223> Composite human delta (H-Delta-1) amino acid
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sequence

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<400> 68
Glu Asn Ser Tyr Ser Cys Thr Cys Pro Pro Gly Phe Tyr Gly Lys Ile
Cys Glu Leu Ser Ala Met Thr Cys Ala Asp Gly Pro Cys Phe Asn Gly
                                25
Gly Arg Cys Ser Asp Ser Pro Asp Gly Gly Tyr Ser Cys Arg Cys Pro
                            40
Val Gly Tyr Ser Gly Phe Asn Cys Glu Lys Lys Ile Asp Tyr Cys Ser
                        55
                                            60
Ser Ser Pro Cys Ser Asn Gly Ala Lys Cys Val Asp Leu Gly Asp Ala
                                        75
Tyr Leu Cys Arg Cys Gln Ala Gly Phe Ser Gly Arg His Cys Asp Asp
                                    90
Asn Val Asp Asp Cys Ala Ser Ser Pro Cys Ala Asn Gly Gly Thr Cys
                                                    110
                                105
Arg Asp Gly Val Asn Asp Phe Ser Cys Thr Cys Pro Pro Gly Tyr Thr
                            120
                                                125
Gly Arg Asn Cys Ser Ala Pro Ala Ser Arg Cys Glu His Ala Pro Cys
                       135
His Asn Gly Ala Thr Cys His Glu Arg Gly His Arg Tyr
145
                    150
<210> 69
<211> 12
<212> PRT
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<223> Composite human delta (H-Delta-1) amino acid
      sequence
<400> 69
Cys Glu Cys Ala Arg Ser Tyr Gly Gly Pro Asn Cys
                 5
<210> 70
<211> 5
<212> PRT
<213> Artificial Sequence
<223> Composite human delta (H-Delta-1) amino acid
      sequence
<400> 70
Phe Leu Leu Pro Glu
<210> 71
<211> 4
<212> PRT
<213> Artificial Sequence
<223> Composite human delta (H-Delta-1) amino acid
      sequence
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<400> 71
Pro Pro Gly Pro
<210> 72
<211> 25
<212> PRT
<213> Artificial Sequence
<220>
<223> Composite human delta (H-Delta-1) amino acid
      sequence
<400> 72
Leu Leu Gly Cys Ala Ala Val Val Cys Val Arg Leu Arg Leu
                                    10
Gln Lys His Arg Pro Pro Ala Asp Pro
            20
                                25
<210> 73
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Composite human delta (H-Delta-1) amino acid
      sequence
<400> 73
Arg Gly Glu Thr Glu Thr Met Asn Asn Leu
                 5
<210> 74
<211> 14
<212> PRT
<213> Artificial Sequence
<220>
<223> Composite human delta (H-Delta-1) amino acid
      sequence
<400> 74
Asn Cys Gln Arg Glu Lys Asp Ile Ser Val Ser Ile Ile Gly
<210> 75
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Composite human delta (H-Delta-1) amino acid
      sequence
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<400> 75
Thr Gln Ile Lys Asn Thr Asn Lys Lys Ala Asp Phe His Gly Asp His
                                    10
<210> 76
<211> 11
<212> PRT
<213> Artificial Sequence
<220>
<223> Composite human delta (H-Delta-1) amino acid
      sequence
<400> 76
Ala Asp Lys Asn Gly Phe Lys Ala Arg Tyr Pro
<210> 77
<211> 26
<212> PRT
<213> Artificial Sequence
<223> Composite human delta (H-Delta-1) amino acid
      sequence
Val Asp Tyr Asn Leu Val Gln Asp Leu Lys Gly Asp Asp Thr Ala Val
Arg Asp Ala His Ser Lys Arg Asp Thr Lys
            20
<210> 78
<211> 13
<212> PRT
<213> Artificial Sequence
<223> Composite human delta (H-Delta-1) amino acid
      sequence
Gln Pro Gln Gly Ser Ser Gly Glu Glu Lys Gly Thr Pro
<210> 79
<211> 4
<212> PRT
<213> Artificial Sequence
<223> Composite human delta (H-Delta-1) amino acid
      sequence
<400> 79
Pro Thr Leu Arg
1
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<210> 80
<211> 4
<212> PRT
<213> Artificial Sequence
<220>
<223> Composite human delta (H-Delta-1) amino acid
      sequence
<400> 80
Arg Lys Arg Pro
<210> 81
<211> 23
<212> DNA
<213> Artificial Sequence
<220>
<223> Degenerated oligo as primer
<220>
<221> VARIANT
<222> 6, 12, 18, 21
\langle 223 \rangle n = I (Inosine)
<400> 81
                                                                       23
ttcggnttya cntggccngg nac
<210> 82
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
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<220>
<221> VARIANT
<222> 3, 9, 12, 15
\langle 223 \rangle n = I (Inosine)
<400> 82
                                                                       20
tcnatgcang tnccnccrtt
<210> 83
<211> 8
<212> PRT
<213> Drosophila
<400> 83
Phe Gly Phe Thr Trp Pro Gly Thr
<210> 84
<211> 7
<212> PRT
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<213> Drosophila
  <400> 84
  Asn Gly Gly Thr Cys Ile Asp
  <210> 85
  <211> 12
  <212> PRT
  <213> Drosophila
<400> 85
  Ser Ile Pro Pro Gly Ser Arg Thr Ser Leu Gly Val
  <210> 86
  <211> 23
  <212> DNA
  <213> Artificial Sequence
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  <223> Primer 1 for PCR
  <220>
  <221> VARIANT
  <222> 3, 9, 15, 18, 21
  \langle 223 \rangle n = I (Inosine)
  <400> 86
  ggnttcacnt ggccnggnac ntt
                                                                         23
  <210> 87
  <211> 23
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  <220>
  <223> Primer 2 for PCR
  <220>
  <221> VARIANT
  <222> 3, 6, 18
  \langle 223 \rangle n = I (Inosine)
  <400> 87
  gtnccnccrt tyttrcangg rtt
                                                                         23
  <210> 88
  <211> 8
  <212> PRT
  <213> Artificial Sequence
  <223> EGF-like repeats encoded by SEQ ID NO. 87
  <400> 88
  Asn Pro Cys Lys Asn Gly Gly Thr
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<210> 89
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<212> DNA
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<223> degenerated oligo primer
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<221> VARIANT
<222> 3, 15, 18
\langle 223 \rangle n = I (Inosine)
<400> 89
acnatgaaya ayctngcnaa ytg
                                                                       23
<210> 90
<211> 8
<212> PRT
<213> Artificial Sequence
<223> amino acid encoded by SEQ ID NO. 89
<400> 90
Thr Met Asn Asn Leu Ala Asn Cys
<210> 91
<211> 23
<212> DNA
<213> Artificial Sequence
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<223> degenerated oligo primer
<220>
<221> VARIANT
<222> 6, 9, 21
\langle 223 \rangle n = I (Inosine)
<400> 91
                                                                       23
acrtanacng aytgrtaytt ngt
<210> 92
<211> 8
<212> PRT
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<220>
<223> amino acid sequence encoded by SEQ ID NO. 91
<400> 92
Thr Lys Tyr Gln Ser Val Tyr Val
<210> 93
<211> 23
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<212> DNA
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<223> degenerated oligo
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<221> VARIANT
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\langle 223 \rangle n = I (Inosine)
<400> 93
gcdatnacrc aytcrtcytt ytc
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<210> 94
<211> 8
<212> PRT
<213> Artificial Sequence
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<223> amino acid sequence endoced by SEQ ID NO. 86
Gly Phe Thr Trp Pro Gly Thr Phe
<210> 95
<211> 129
<212> PRT
<213> Homo sapiens
<400> 95
Thr Met Asn Asn Leu Ala Asn Cys Gln Arg Glu Lys Asp Ile Ser Ile
                                     10
Ser Val Ile Gly Ala Thr Gln Ile Lys Asn Thr Asn Lys Lys Val Asp
                                25
Phe His Ser Asp Asn Ser Asp Lys Asn Gly Tyr Lys Val Arg Tyr Pro
                            40
Ser Val Asp Tyr Asn Leu Val His Glu Leu Lys Asn Glu Asp Ser Val
                        55
Lys Glu Glu His Gly Lys Cys Glu Ala Lys Cys Glu Thr Tyr Asp Ser
                    70
Glu Ala Glu Glu Lys Ser Ala Val Gln Leu Lys Ser Ser Asp Thr Ser
                                     90
Glu Arg Lys Arg Pro Asp Ser Val Tyr Ser Thr Ser Lys Asp Thr Lys
            100
                                105
Tyr Gln Ser Val Tyr Val Ile Ser Glu Glu Lys Asp Glu Cys Ile Ile
       115
                            120
                                                 125
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Ala